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<212> PRT

<213> Streptococcus agalactiae

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Thr	Val	Thr 35	Glu	Asp	Thr	Pro	Ala 40	Thr	Glu	Gln	Ala	Val 45	Glu	Pro	Pro
Gln	Pro 50	Ile	Ala	Val	Ser	Glu 55	Glu	Ser	Arg	Ser	Ser 60	Lys	Glu	Thr	Lys
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Asn	Asp	Leu	Ala	Pro 85	Gln	Ala	Pro	Ala	Lys 90	Thr	Ala	Asp	Thr	Pro 95	Ala
Thr	Ser	Lys	Ala 100	Thr	Ile	Arg	Asp	Leu 105	Asn	Asp	Pro	Ser	His 110	Val	Lys
Thr	Leu	Gln 115	Glu	Lys	Ala	Gly	Lys 120	Gly	Ala	Gly	Thr	Val 125	Val	Ala	Val
Ile	Asp 130	Ala	Gly	Phe	Asp	Lys 135	Asn	His	Glu	Ala	Trp 140	Arg	Leu	Thr	Asp
Lys 145	Thr	Lys	Ala	Arg	Tyr 150	Gln	Ser	Lys	Glu	Asn 155	Leu	Glu	Lys	Ala	Lys 160
Lys	Glu	His	Gly	Ile 165	Thr	Tyr	Gly	Glu	Trp 170	Val	Asn	Asp	Lys	Val 175	Ala
Tyr	Tyr	His	Asp 180	Tyr	Ser	Lys	Asp	Gly 185	Lys	Asn	Ala	Val	Asp 190	Gln	Glu
His	Gly	Thr 195	His	Val	Ser	Gly	Ile 200	Leu	Ser	Gly	Asn	Ala 205	Pro	Ser	Glu
Met	Lys 210	Glu	Pro	Tyr	Arg	Leu 215	Glu	Gly	Ala	Met	Pro 220	Glu	Ala	Gln	Leu
Leu 225	Leu	Met	Arg	Val	Glu 230	Ile	Val	Asn	Gly	Leu 235	Ala	Asp	Tyr	Ala	Arg 240
Asn	Tyr	Ala	Gln	Ala	Ile	Arg	Asp	Ala	Val	Asn	Leu	Gly	Ala	Lys	Val

Ile Asn Met Ser Phe Gly Asn Ala Ala Leu Ala Tyr Ala Asn Leu Pro Asp Glu Thr Lys Lys Ala Phe Asp Tyr Ala Lys Ser Lys Gly Val Ser Ile Val Thr Ser Ala Gly Asn Asp Ser Ser Phe Gly Gly Lys Pro Arg Leu Pro Leu Ala Asp His Pro Asp Tyr Gly Val Val Gly Thr Pro Ala Ala Ala Asp Ser Thr Leu Thr Val Ala Ser Tyr Ser Pro Asp Lys Gln Leu Thr Glu Thr Ala Thr Val Lys Thr Asp Asp His Gln Asp Lys Glu Met Pro Val Ile Ser Thr Asn Arg Phe Glu Pro Asn Lys Ala Tyr Asp Tyr Ala Tyr Ala Asn Arg Gly Thr Lys Glu Asp Asp Phe Lys Asp Val Glu Gly Lys Ile Ala Leu Ile Glu Arg Gly Asp Ile Asp Phe Lys Asp Lys Ile Ala Asn Ala Lys Lys Ala Gly Ala Val Gly Val Leu Ile Tyr Asp Asn Gln Asp Lys Gly Phe Pro Ile Glu Leu Pro Asn Val Asp Gln Met Pro Ala Ala Phe Ile Ser Arg Arg Asp Gly Leu Leu Leu Lys Asp Asn Pro Pro Lys Thr Ile Thr Phe Asn Ala Thr Pro Lys Val Leu Pro Thr Ala Ser Gly Thr Lys Leu Ser Arg Phe Ser Ser Trp Gly Leu Thr Ala Asp Gly Asn Ile Lys Pro Asp Ile Ala Ala Pro Gly Gln Asp Ile Leu Ser Ser Val Ala Asn Asn Lys Tyr Ala Lys Leu Ser Gly Thr Ser Met Ser Ala Pro Leu Val Ala Gly Ile Met Gly Leu Leu Gln Lys Gln Tyr Glu Thr Gln Tyr Pro Asp Met Thr Pro Ser Glu Arg Leu Asp Leu Ala Lys Lys Val Leu Met Ser Ser Ala Thr Ala Leu Tyr Asp Glu Asp Glu Lys Ala Tyr Phe Ser Pro Arg Gln Gln Gly Ala Gly Ala Val Asp Ala Lys Lys Ala Ser Ala Ala Thr Met Tyr Val Thr Asp Lys Asp Asn Thr Ser Ser Lys Val His Leu Asn Asn Val Ser Asp Lys Phe Glu Val Thr Val Thr Val His Asn Lys Ser Asp Lys Pro Gln Glu Leu Tyr Tyr Gln Val Thr Val Gln Thr Asp Lys Val Asp Gly Lys His Phe Ala Leu Ala Pro Lys Ala Leu Tyr Glu Thr Ser Trp Gln Lys Ile Thr Ile Pro Ala Asn Ser Ser Lys Gln Val Thr Val Pro Ile Asp Ala Ser Arg Phe Ser Lys Asp Leu Leu Ala Gln Met Lys Asn Gly Tyr Phe Leu Glu Gly

Phe Val Arg Phe Lys Gln Asp Pro Thr Lys Glu Glu Leu Met Ser Ile Pro Tyr Ile Gly Phe Arg Gly Asp Phe Gly Asn Leu Ser Ala Leu Glu Lys Pro Ile Tyr Asp Ser Lys Asp Gly Ser Ser Tyr Tyr His Glu Ala Asn Ser Asp Ala Lys Asp Gln Leu Asp Gly Asp Gly Leu Gln Phe Tyr Ala Leu Lys Asn Asn Phe Thr Ala Leu Thr Thr Glu Ser Asn Pro Trp Thr Ile Ile Lys Ala Val Lys Glu Gly Val Glu Asn Ile Glu Asp Ile Glu Ser Ser Glu Ile Thr Glu Thr Ile Phe Ala Gly Thr Phe Ala Lys Gln Asp Asp Ser His Tyr Tyr Ile His Arg His Ala Asn Gly Lys Pro Tyr Ala Ala Ile Ser Pro Asn Gly Asp Gly Asn Arg Asp Tyr Val Gln Phe Gln Gly Thr Phe Leu Arg Asn Ala Lys Asn Leu Val Ala Glu Val Leu Asp Lys Glu Gly Asn Val Val Trp Thr Ser Glu Val Thr Glu Gln Val Val Lys Asn Tyr Asn Asn Asp Leu Ala Ser Thr Leu Gly Ser Thr Arg Phe Glu Lys Thr Arg Trp Asp Gly Lys Asp Lys Asp Gly Lys Val Val Ala Asn Gly Thr Tyr Thr Tyr Arg Val Arg Tyr Thr Pro Ile Ser Ser Gly Ala Lys Glu Gln His Thr Asp Phe Asp Val Ile Val Asp Asn Thr Thr Pro Glu Val Ala Thr Ser Ala Thr Phe Ser Thr Glu Asp Ser Arg Leu Thr Leu Ala Ser Lys Pro Lys Thr Ser Gln Pro Val Tyr Arg Glu Arg Ile Ala Tyr Thr Tyr Met Asp Glu Asp Leu Pro Thr Thr Glu Tyr Ile Ser Pro Asn Glu Asp Gly Thr Phe Thr Leu Pro Glu Glu Ala Glu Thr Met Glu Gly Ala Thr Val Pro Leu Lys Met Ser Asp Phe Thr Tyr Val Val Glu Asp Met Ala Gly Asn Ile Thr Tyr Thr Pro Val Thr Lys Leu Leu Glu Gly His Ser Asn Lys Pro Glu Gln Asp Gly Ser Asp Gln Ala Pro Asp Lys Lys Pro Glu Ala Lys Pro Glu Gln Asp Gly Ser Gly Gln Thr Pro Asp Lys Lys Glu Thr Lys Pro Glu Lys Asp Ser Ser Gly Gln Thr Pro Gly Lys Thr Pro Gln Lys Gly Gln Ser Ser Arg Thr Leu Glu Lys Arg Ser Ser Lys Arg Ala Leu Ala Thr Lys Ala Ser Thr Arg Asp Gln Leu Pro Thr Thr Asn Asp Lys Asp Thr Asn Arg Leu His Leu Leu Lys Leu Val Met Thr Thr Phe Phe Leu Gly

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attaaagtgg tttgtggtag tcatccttta gaattgttag atgaggattt ttgttacatg
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gctgcgaatg ataaagatac tctagttatg gaattatcaa gttttcagct aatgggagtt
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aatcaagtta ttagagaaac tttaagcaat tttggaggtg ttaaacaccg cttgcaatca
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                                                                      1080
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gttttagggg aatcggcatc tcgagtaaaa cgtgctgcac aaaaagcagg agtaacttat
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                            40
Glu Asn Pro Thr Ala Gln Ser Leu Leu Glu Glu Gly Ile Lys Val Val
                        55
Cys Gly Ser His Pro Leu Glu Leu Leu Asp Glu Asp Phe Cys Tyr Met
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                                        75
Ile Lys Asn Pro Gly Ile Pro Tyr Asn Asn Pro Met Val Lys Lys Ala
                                    90
Leu Glu Lys Gln Ile Pro Val Leu Thr Glu Val Glu Leu Ala Tyr Leu
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Val Ser Glu Ser Gln Leu Ile Gly Ile Thr Gly Ser Asn Gly Lys Thr
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120

180

420 480

540

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Leu Met Gly Val Lys Glu Phe Arg Pro His Ile Ala Val Ile Thr Asn
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Leu Met Pro Thr His Leu Asp Tyr His Gly Ser Phe Glu Asp Tyr Val
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                                               205
Ala Ala Lys Trp Asn Ile Gln Asn Gln Met Ser Ser Asp Phe Leu
                       215
                                           220
Val Leu Asn Phe Asn Gln Gly Ile Ser Lys Glu Leu Ala Lys Thr Thr
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                                       235
Lys Ala Thr Ile Val Pro Phe Ser Thr Thr Glu Lys Val Asp Gly Ala
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Tyr Val Gln Asp Lys Gln Leu Phe Tyr Lys Gly Glu Asn Ile Met Ser
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Val Asp Asp Ile Gly Val Pro Gly Ser His Asn Val Glu Asn Ala Leu
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Ala Thr Ile Ala Val Ala Lys Leu Ala Gly Ile Ser Asn Gln Val Ile
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Arg Glu Thr Leu Ser Asn Phe Gly Gly Val Lys His Arg Leu Gln Ser
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                                       315
Leu Gly Lys Val His Gly Ile Ser Phe Tyr Asn Asp Ser Lys Ser Thr
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Asn Ile Leu Ala Thr Gln Lys Ala Leu Ser Gly Phe Asp Asn Thr Lys
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Val Ile Leu Ile Ala Gly Gly Leu Asp Arg Gly Asn Glu Phe Asp Glu
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                                               365
Leu Ile Pro Asp Ile Thr Gly Leu Lys His Met Val Val Leu Gly Glu
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                                           380
Ser Ala Ser Arg Val Lys Arg Ala Ala Gln Lys Ala Gly Val Thr Tyr
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                                       395
Ser Asp Ala Leu Asp Val Arg Asp Ala Val His Lys Ala Tyr Glu Val
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                                   410
Ala Gln Gln Gly Asp Val Ile Leu Leu Ser Pro Ala Asn Ala Ser Trp
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                                                                     420
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Glu Val Tyr Gly Ile Asn Gln Gly Tyr Tyr Gly Met Val Thr Gly Asp
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Ile Phe Pro Leu Asp Ala Asn Ser Val Gly Asp Thr Ile Asn Arg Gly
                        55
                                            60
Gly Thr Phe Leu Arg Ser Ala Arg Tyr Pro Glu Phe Ala Glu Leu Glu
                    70
                                        75
Gly Gln Leu Lys Gly Ile Glu Gln Leu Lys Lys His Gly Ile Glu Gly
                85
                                    90
Val Val Ile Gly Gly Asp Gly Ser Tyr His Gly Ala Met Arg Leu
                                105
Thr Glu His Gly Phe Pro Ala Val Gly Leu Pro Gly Thr Ile Asp Asn
                            120
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Asp Ile Val Gly Thr Asp Tyr Thr Ile Gly Phe Asp Thr Ala Val Ala
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                                            140
Thr Ala Val Glu Asn Leu Asp Arg Leu Arg Asp Thr Ser Ala Ser His
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                                        155
Asn Arg Thr Phe Val Val Glu Val Met Gly Arg Asn Ala Gly Asp Ile
                                    170
                165
                                                        175
Ala Leu Trp Ser Gly Ile Ala Ala Gly Ala Asp Gln Ile Ile Val Pro
                                185
Glu Glu Glu Phe Asn Ile Asp Glu Val Val Ser Asn Val Arg Ala Gly
                            200
                                                205
Tyr Ala Ala Gly Lys His His Gln Ile Ile Val Leu Ala Glu Gly Val
Met Ser Gly Asp Glu Phe Ala Lys Thr Met Lys Ala Ala Gly Asp Asp
                    230
                                        235
Ser Asp Leu Arg Val Thr Asn Leu Gly His Leu Leu Arg Gly Gly Ser
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Pro Thr Ala Arg Asp Arg Val Leu Ala Ser Arg Met Gly Ala Tyr Ala
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                                265
Val Gln Leu Leu Lys Glu Gly Arg Gly Gly Leu Ala Val Gly Val His
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                                                285
Asn Glu Glu Met Val Glu Ser Pro Ile Leu Gly Leu Ala Glu Gly
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600

660

720

780

840

900 960

1020

300

295

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Gln Ser Ser Lys
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                                                                       180
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Ser Ser Tyr Thr Val Lys Tyr Gly Asp Thr Leu Ser Val Ile Ser Glu
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Ala Met Ser Ile Asp Met Asn Val Leu Ala Lys Ile Asn Asn Ile Ala
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Asp Ile Asn Leu Ile Tyr Pro Glu Thr Thr Leu Thr Val Thr Tyr Asp
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Asn Ala Ala Gly Gln Thr Thr Ala Thr Val Asp Leu Lys Thr Asn Gln
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Val Ser Val Ala Asp Gln Lys Val Ser Leu Asn Thr Ile Ser Glu Gly
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Met Thr Pro Glu Ala Ala Thr Thr Ile Val Ser Pro Met Lys Thr Tyr
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Ser Ser Ala Pro Ala Leu Lys Ser Lys Glu Val Leu Ala Gln Glu Gln
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Ala Val Ser Gln Ala Ala Ala Asn Glu Gln Val Ser Pro Ala Pro Val
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Lys Ser Ile Thr Ser Glu Val Pro Ala Ala Lys Glu Glu Val Lys Pro
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Thr Gln Thr Ser Val Ser Gln Ser Thr Thr Val Ser Pro Ala Ser Val
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Ala Ala Glu Thr Pro Ala Pro Val Ala Lys Val Ala Pro Val Arg Thr
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Val Ala Ala Pro Arg Val Ala Ser Val Lys Val Val Thr Pro Lys Val
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Glu Thr Gly Ala Ser Pro Glu His Val Ser Ala Pro Ala Val Pro Val
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Thr Thr Thr Ser Pro Ala Thr Asp Ser Lys Leu Gln Ala Thr Glu Val
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Lys Ser Val Pro Val Ala Gln Lys Ala Pro Thr Ala Thr Pro Val Ala
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                                             300
Gln Pro Ala Ser Thr Thr Asn Ala Val Ala Ala His Pro Glu Asn Ala
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                                        315
Gly Leu Gln Pro His Val Ala Ala Tyr Lys Glu Lys Val Ala Ser Thr
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Asn Ile Ser Tyr Val Ile Trp Gln Gln Lys Phe Tyr Ser Asn Thr Asn
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60

120

180 240

300

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Gln Leu Asp Ala Tyr Met Asp Asp Ala Gln Lys Asp Phe Lys Gln Thr
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Asn Phe Asn Ala Met Asn Val Glu Tyr Gly Thr Leu Gly Asn His Glu
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<213> Streptococcus agalactiae

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Thr Ile His Lys Lys Gln Leu His Gln Phe Thr Ala Ile Asn Pro Met
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                    630
Arg Asn Tyr Gly Lys Pro Ser Asn Ser Thr Thr Val Lys Ser Lys Gln
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Leu Pro Lys Thr Asn Ser Glu Tyr Gly Gln Ser Phe Leu Met Ser Val
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Val Ala Leu Glu Ala Gly Phe Val Glu Ser Gly Asp Asn Ile Val Ile
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Gln Ala Leu Leu Lys Pro Ser Lys Lys Val Phe Gln Ile Ile Asp His
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Ile Gln Leu Ala Leu Lys Pro Val Asn Val Arg Phe Gly Leu Gly Thr
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Gly Asn Ile Ile Thr Ser Ile Asn Ser Asn Glu Ser Ile Gly Ala Asp
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Val Met Lys Ala Ser Val Ile Asp Gly Leu Thr Pro Ser Lys Ser Ile
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Ser Gly Leu Thr Gly Gly Asp Ala Val Lys Met Asp Gln Tyr Leu Gln
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Ser Gly Lys Thr Ile Ser Asp Thr Thr Ile Leu Ala Ala Val Arg Asn
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Phe Thr Ala Gly Ala Phe Gly Leu Val Ile Gly Asn Asn Ala Ser Ile
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Ala Ser Gln Ala Ile Ala Phe Val Ile Lys Asn Met Leu Gly Leu Ile
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Cys Asp Pro Val Ala Gly Leu Val Glu Val Pro Cys Val Lys Arg Asn
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Gly Ile Glu Ser Gln Ile Pro Val Asp Glu Val Ile Asp Ala Met Tyr
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110

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1500

1560

1620

1680

1740

1800

1860

1920

1980

2040

2100

2160

2193

265

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                                                                      360
aaacttgttc tttacaccaa ttcaattcaa acaggttgtg gtataggtga atctgcttca
                                                                      420
                                                                      480
ggaccatttt attgttcagc agataaaaaa atctatcttg atatttcttt ttacaatgaa
                                                                      540
ttatcacata aatatggtgc tactggtgat tttgctatgg cctacgtcat cgcccacgaa
                                                                      600
gttggtcacc acattcaaac agagttaggc attatggata agtataatag aatgcgacac
ggacttacta agaaagaagc aaatgcttta aatgttcggc tagaacttca agcagattat
                                                                      660
                                                                      720
tatgcagggg tatgggctca ctacatcagg ggaaaaaatc tcttagaaca aggagacttt
gaagaggcca tgaatgctgc ccacgccgtc ggagacgata cccttcagaa agaaacctac
                                                                      780
                                                                      840
ggaaaattag tgcctgatag ctttacccat ggaacagctg aacaacgcca acgttggttt
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<210> 30
<211> 300
<212> PRT
<213> Streptococcus agalactiae
<400> 30
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Ser Ser Ser Gly Gly Ser Phe Ser Ser Gly Gly Ser Gly Leu Pro Ile
                                25
Leu Gln Leu Leu Leu Arg Gly Ser Trp Lys Thr Lys Leu Val Val
                            40
                                                45
Leu Ile Ile Leu Leu Leu Gly Gly Gly Leu Thr Ser Ile Phe
                        55
                                            60
Asn Asp Ser Ser Ser Pro Ser Ser Tyr Gln Ser Gln Asn Val Ser Arg
                    70
                                        75
Ser Val Asp Asn Ser Ala Thr Arg Glu Gln Ile Asp Phe Val Asn Lys
Val Leu Gly Ser Thr Glu Asp Phe Trp Ser Gln Glu Phe Gln Thr Gln
                                105
                                                    110
Gly Phe Gly Asn Tyr Lys Glu Pro Lys Leu Val Leu Tyr Thr Asn Ser
                            120
Ile Gln Thr Gly Cys Gly Ile Gly Glu Ser Ala Ser Gly Pro Phe Tyr
                        135
                                            140
Cys Ser Ala Asp Lys Lys Ile Tyr Leu Asp Ile Ser Phe Tyr Asn Glu
                    150
                                        155
Leu Ser His Lys Tyr Gly Ala Thr Gly Asp Phe Ala Met Ala Tyr Val
                165
                                    170
Ile Ala His Glu Val Gly His His Ile Gln Thr Glu Leu Gly Ile Met
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```
185
Asp Lys Tyr Asn Arg Met Arg His Gly Leu Thr Lys Lys Glu Ala Asn
                            200
Ala Leu Asn Val Arg Leu Glu Leu Gln Ala Asp Tyr Tyr Ala Gly Val
                        215
                                            220
Trp Ala His Tyr Ile Arg Gly Lys Asn Leu Leu Glu Gln Gly Asp Phe
                    230
                                        235
Glu Glu Ala Met Asn Ala Ala His Ala Val Gly Asp Asp Thr Leu Gln
                245
                                    250
Lys Glu Thr Tyr Gly Lys Leu Val Pro Asp Ser Phe Thr His Gly Thr
                                265
Ala Glu Gln Arg Gln Arg Trp Phe Asn Lys Gly Phe Gln Tyr Gly Asp
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Ile Gln His Gly Asp Thr Phe Ser Val Glu His Leu
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<212> DNA
<213> Streptococcus agalactiae
<400> 31
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                                                                       120
aattcatctg ttgatacgag ccaggaattt caaaataatt taaaaaatgc tattggtaac
                                                                       180
ctaccatttc aatatgttaa tggtatttat gaattaaata ataatcagac aaatttaaat
                                                                       240
gctgatgtca atgttaaagc gtatgttcaa aatacaattg acaatcaaca aagactatca
                                                                       300
actgctaatg caatgcttga tagaaccatt cgtcaatatc aaaatcgcag agataccact
                                                                       360
cttcccgatg caaattggaa accattaggt tggcatcaag tagctactaa tgaccattat
                                                                       420
ggacatgcag tcgacaaggg gcatttaatt gcctatgctt tagctggaaa tttcaaaggt
                                                                       480
tgggatgctt ccgtgtcaaa tcctcaaaat gttgtcacac aaacagctca ttccaaccaa
                                                                       540
tcaaatcaaa aaatcaatcg tggacaaaat tattatgaaa gcttagttcg taaggcggtt
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gaccaaaaca aacgtgttcg ttaccgtgta actccattgt accgtaatga tactgattta
                                                                       660
gttccatttg caatgcacct agaagctaaa tcacaagatg gcacattaga atttaatgtt
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                                                                       783
aat
<210> 32
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<212> PRT
<213> Streptococcus agalactiae
<400> 32
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Met Leu Gly Val Met Thr Phe Gly Leu Pro Thr Gln Pro Gln Asn Val
                                25
                                                     30
Thr Pro Ile Val His Ala Asp Val Asn Ser Ser Val Asp Thr Ser Gln
                            40
Glu Phe Gln Asn Asn Leu Lys Asn Ala Ile Gly Asn Leu Pro Phe Gln
                        55
Tyr Val Asn Gly Ile Tyr Glu Leu Asn Asn Asn Gln Thr Asn Leu Asn
                    70
                                        75
Ala Asp Val Asn Val Lys Ala Tyr Val Gln Asn Thr Ile Asp Asn Gln
                85
                                    90
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Gln Arg Leu Ser Thr Ala Asn Ala Met Leu Asp Arg Thr Ile Arg Gln
                                105
Tyr Gln Asn Arg Arg Asp Thr Thr Leu Pro Asp Ala Asn Trp Lys Pro
                            120
Leu Gly Trp His Gln Val Ala Thr Asn Asp His Tyr Gly His Ala Val
                        135
                                            140
Asp Lys Gly His Leu Ile Ala Tyr Ala Leu Ala Gly Asn Phe Lys Gly
                    150
                                        155
Trp Asp Ala Ser Val Ser Asn Pro Gln Asn Val Val Thr Gln Thr Ala
                165
                                    170
His Ser Asn Gln Ser Asn Gln Lys Ile Asn Arg Gly Gln Asn Tyr Tyr
                                185
Glu Ser Leu Val Arg Lys Ala Val Asp Gln Asn Lys Arg Val Arg Tyr
                            200
                                                205
Arg Val Thr Pro Leu Tyr Arg Asn Asp Thr Asp Leu Val Pro Phe Ala
                        215
                                            220
Met His Leu Glu Ala Lys Ser Gln Asp Gly Thr Leu Glu Phe Asn Val
                    230
                                        235
Ala Ile Pro Asn Thr Gln Ala Ser Tyr Thr Met Asp Tyr Ala Thr Gly
                245
                                    250
Glu Ile Thr Leu Asn
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<211> 1242
<212> DNA
<213> Streptococcus agalactiae
<400> 33
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                                                                       120
                                                                       180
gcagtaaaaa ctaactacaa agtttttaat gttagagaag gaagtgtttc gtcctcaact
                                                                       240
cttttgacag gaaaagctaa ggctaatcaa gaacagtatg tgtattttga tgctaataaa
                                                                       300
ggtaatcgag caactgtcac agttaaagtg ggtgataaaa tcacagctgg tcagcagtta
gttcaatatg atacaacaac tgcacaagca gcctacgaca ctgctaatcg tcaattaaat
                                                                       360
                                                                       420
aaagtagcgc gtcagattaa taatctaaag acaacaggaa gtcttccagc tatggaatca
                                                                       480
agtgatcaat cttcttcatc atcacaagga caagggactc aatcgactag tggtgcgacg
aatcgtctac agcaaaatta tcaaagtcaa gctaatgctt catacaacca acaacttcaa
                                                                       540
gatttgaatg atgcttatgc agatgcacag gcagaagtaa ataaagcaca aaaagcattg
                                                                       600
                                                                       660
aatgatactg ttattacaag tgacgtatca gggacagttg ttgaagttaa tagtgatatt
                                                                       720
gatccagctt caaaaactag tcaagtactt gtccatgtag caactgaagg taaactccaa
                                                                       780
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ataaaatcta aggtctatcc tgacaaggaa tgggaaggta aaatttcata tatctcaaat
                                                                       840
tatccagaag cagaagcaaa caacaatgac tctaataacg gctctagtgc tgtaaattat
                                                                       900
aaatataaag tagatattac tagccctctc gatgcattaa aacaaggttt taccgtatca
                                                                       960
gttgaagtag ttaatggaga taagcacctt attgtcccta caagttctgt gataaacaaa
                                                                      1020
gataataaac actttgtttg ggtatacaat gattctaatc gtaaaatttc caaagttgaa
                                                                      1080
gtcaaaattg gtaaagctga tgctaagaca caagaaattt tatcaggttt gaaagcagga
                                                                      1140
caaatcgtgg ttactaatcc aagtaaaacc ttcaaggatg ggcaaaaaaat tgataatatt
                                                                      1200
gaatcaatcg atcttaactc taataagaaa tcagaggtga aa
                                                                      1242
<210> 34
<211> 414
<212> PRT
<213> Streptococcus agalactiae
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<400> 34 Met Ser Lys Arg Gln Asn Leu Gly Ile Ser Lys Lys Gly Ala Ile Ile 10 Ser Gly Leu Ser Val Ala Leu Ile Val Val Ile Gly Gly Phe Leu Trp 25 Val Gln Ser Gln Pro Asn Lys Ser Ala Val Lys Thr Asn Tyr Lys Val Phe Asn Val Arg Glu Gly Ser Val Ser Ser Ser Thr Leu Leu Thr Gly Lys Ala Lys Ala Asn Gln Glu Gln Tyr Val Tyr Phe Asp Ala Asn Lys 70 75 Gly Asn Arg Ala Thr Val Thr Val Lys Val Gly Asp Lys Ile Thr Ala 90 Gly Gln Gln Leu Val Gln Tyr Asp Thr Thr Thr Ala Gln Ala Ala Tyr 105 Asp Thr Ala Asn Arg Gln Leu Asn Lys Val Ala Arg Gln Ile Asn Asn 120 Leu Lys Thr Thr Gly Ser Leu Pro Ala Met Glu Ser Ser Asp Gln Ser 135 140 Ser Ser Ser Gln Gly Gln Gly Thr Gln Ser Thr Ser Gly Ala Thr 150 155 Asn Arg Leu Gln Gln Asn Tyr Gln Ser Gln Ala Asn Ala Ser Tyr Asn 165 170 Gln Gln Leu Gln Asp Leu Asn Asp Ala Tyr Ala Asp Ala Gln Ala Glu 180 185 Val Asn Lys Ala Gln Lys Ala Leu Asn Asp Thr Val Ile Thr Ser Asp 200 Val Ser Gly Thr Val Val Glu Val Asn Ser Asp Ile Asp Pro Ala Ser 215 220 Lys Thr Ser Gln Val Leu Val His Val Ala Thr Glu Gly Lys Leu Gln 230 235 Val Gln Gly Thr Met Ser Glu Tyr Asp Leu Ala Asn Val Lys Lys Asp 245 250 Gln Ala Val Lys Ile Lys Ser Lys Val Tyr Pro Asp Lys Glu Trp Glu 265 Gly Lys Ile Ser Tyr Ile Ser Asn Tyr Pro Glu Ala Glu Ala Asn Asn 280 285 Asn Asp Ser Asn Asn Gly Ser Ser Ala Val Asn Tyr Lys Tyr Lys Val 295 300 Asp Ile Thr Ser Pro Leu Asp Ala Leu Lys Gln Gly Phe Thr Val Ser 310 315 Val Glu Val Val Asn Gly Asp Lys His Leu Ile Val Pro Thr Ser Ser 325 330 Val Ile Asn Lys Asp Asn Lys His Phe Val Trp Val Tyr Asn Asp Ser 345 Asn Arg Lys Ile Ser Lys Val Glu Val Lys Ile Gly Lys Ala Asp Ala 360 Lys Thr Gln Glu Ile Leu Ser Gly Leu Lys Ala Gly Gln Ile Val Val 375 380 Thr Asn Pro Ser Lys Thr Phe Lys Asp Gly Gln Lys Ile Asp Asn Ile 390 395 Glu Ser Ile Asp Leu Asn Ser Asn Lys Lys Ser Glu Val Lys 405 410

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<211> 930
<212> DNA
<213> Streptococcus agalactiae
<400> 35
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caacaaacta aacaagaaag cactaaaaca actatttcta aaatgcctaa aattgaaggc
ttcacctatt atggaaaaat tcctgaaaat ccgaaaaaag taattaattt tacatattct
tacactgggt atttattaaa actaggtgtt aatgtttcaa gttacagttt agacttagaa
aaagatagcc ccgtttttgg taaacaactg aaagaagcta aaaaattaac tgctgatgat
acagaagcta ttgccgcaca aaaacctgat ttaatcatgg ttttcgatca agatccaaac
atcaatactc tgaaaaaaat tgcaccaact ttagttatta aatatggtgc acaaaattat
ttagatatga tgccagcctt ggggaaagta ttcggtaaag aaaaagaagc taatcagtgg
gttagccaat ggaaaactaa aactctcgct gtcaaaaaaag atttacacca tatcttaaag
cctaacacta cttttactat tatggatttt tatgataaaa atatctattt atatggtaat
aattttggac gcggtggaga actaatctat gattcactag gttatgctgc cccagaaaaa
gtcaaaaaag atgtctttaa aaaagggtgg tttaccgttt cgcaagaagc aatcggtgat
tacgttggag attatgccct tgttaatata aacaaaacga ctaaaaaagc agcttcatca
cttaaagaaa gtgatgtctg gaagaattta ccagctgtca aaaaagggca catcatagaa
agtaactacg acgtgtttta tttctctgac cctctatctt tagaagctca attaaaatca
tttacaaagg ctatcaaaga aaatacaaat
<210> 36
<211> 310
<212> PRT
<213> Streptococcus agalactiae
<400> 36
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Val Ser Cys Gly Gln Gln Thr Lys Gln Glu Ser Thr Lys Thr Thr Ile
                                25
Ser Lys Met Pro Lys Ile Glu Gly Phe Thr Tyr Tyr Gly Lys Ile Pro
                            40
Glu Asn Pro Lys Lys Val Ile Asn Phe Thr Tyr Ser Tyr Thr Gly Tyr
                        55
                                            60
Leu Leu Lys Leu Gly Val Asn Val Ser Ser Tyr Ser Leu Asp Leu Glu
                                        75
                    70
Lys Asp Ser Pro Val Phe Gly Lys Gln Leu Lys Glu Ala Lys Lys Leu
                85
                                    90
                                                        95
Thr Ala Asp Asp Thr Glu Ala Ile Ala Ala Gln Lys Pro Asp Leu Ile
                                105
Met Val Phe Asp Gln Asp Pro Asn Ile Asn Thr Leu Lys Lys Ile Ala
                            120
                                                125
Pro Thr Leu Val Ile Lys Tyr Gly Ala Gln Asn Tyr Leu Asp Met Met
                        135
                                            140
Pro Ala Leu Gly Lys Val Phe Gly Lys Glu Lys Glu Ala Asn Gln Trp
                                        155
Val Ser Gln Trp Lys Thr Lys Thr Leu Ala Val Lys Lys Asp Leu His
                                    170
                165
His Ile Leu Lys Pro Asn Thr Thr Phe Thr Ile Met Asp Phe Tyr Asp
                                185
                                                    190
Lys Asn Ile Tyr Leu Tyr Gly Asn Asn Phe Gly Arg Gly Glu Leu
        195
                            200
                                                205
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120

180

240

300

360

420

480

540 600

660

720

780 840

900

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Ile Tyr Asp Ser Leu Gly Tyr Ala Ala Pro Glu Lys Val Lys Lys Asp
    210
                        215
                                            220
Val Phe Lys Lys Gly Trp Phe Thr Val Ser Gln Glu Ala Ile Gly Asp
                    230
                                        235
Tyr Val Gly Asp Tyr Ala Leu Val Asn Ile Asn Lys Thr Thr Lys Lys
                245
                                    250
Ala Ala Ser Ser Leu Lys Glu Ser Asp Val Trp Lys Asn Leu Pro Ala
                                265
Val Lys Lys Gly His Ile Ile Glu Ser Asn Tyr Asp Val Phe Tyr Phe
                            280
Ser Asp Pro Leu Ser Leu Glu Ala Gln Leu Lys Ser Phe Thr Lys Ala
                        295
                                            300
Ile Lys Glu Asn Thr Asn
<210> 37
<211> 576
<212> DNA
<213> Streptococcus agalactiae
<400> 37
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tattcatcaa teggttatge tgatacaagt gataagaata etgacaegag tgtegtgaet
                                                                       120
acgaccttat ctgaggagaa aagatcagat gaactagacc agtctagtac tggttcttct
                                                                       180
tctgaaaatg aatcgagttc atcaagtgaa ccagaaacaa atccgtcaac taatccacct
                                                                       240
acaacagaac catcgcaacc ctcacctagt gaagagaaca agcctgatgg tagaacgaag
                                                                       300
acagaaattg gcaataataa ggatatttct agtggaacaa aagtattaat ttcagaagat
                                                                       360
agtattaaga attttagtaa agcaagtagt gatcaagaag aagtggatcg cgatgaatca
                                                                       420
tcatcttcaa aagcaaatga tgggaaaaaa ggccacagta agcctaaaaa ggaacttcct
                                                                       480
aaaacaggag atagccactc agatactgta atagcatcta cgggagggat tattctgtta
                                                                       540
tcattaagtt tttacaataa gaaaatgaaa ctttat
                                                                       576
<210> 38
<211> 192
<212> PRT
<213> Streptococcus agalactiae
<400> 38
Met Lys Val Lys Asn Lys Ile Leu Thr Met Val Ala Leu Thr Val Leu
                 5
                                    10
Thr Cys Ala Thr Tyr Ser Ser Ile Gly Tyr Ala Asp Thr Ser Asp Lys
                                25
Asn Thr Asp Thr Ser Val Val Thr Thr Thr Leu Ser Glu Glu Lys Arg
                            40
Ser Asp Glu Leu Asp Gln Ser Ser Thr Gly Ser Ser Ser Glu Asn Glu
                        55
                                            60
Ser Ser Ser Ser Glu Pro Glu Thr Asn Pro Ser Thr Asn Pro Pro
                    70
                                        75
Thr Thr Glu Pro Ser Gln Pro Ser Pro Ser Glu Glu Asn Lys Pro Asp
                                    90
Gly Arg Thr Lys Thr Glu Ile Gly Asn Asn Lys Asp Ile Ser Ser Gly
                                105
Thr Lys Val Leu Ile Ser Glu Asp Ser Ile Lys Asn Phe Ser Lys Ala
                            120
        115
Ser Ser Asp Gln Glu Glu Val Asp Arg Asp Glu Ser Ser Ser Lys
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130
                        135
                                            140
Ala Asn Asp Gly Lys Lys Gly His Ser Lys Pro Lys Lys Glu Leu Pro
                    150
                                        155
Lys Thr Gly Asp Ser His Ser Asp Thr Val Ile Ala Ser Thr Gly Gly
                165
                                    170
                                                         175
Ile Ile Leu Leu Ser Leu Ser Phe Tyr Asn Lys Lys Met Lys Leu Tyr
            180
                                185
<210> 39
<211> 924
<212> DNA
<213> Streptococcus agalactiae
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                                                                       120
ccagtatatt ccattacaaa agcagtttct ggtgatttga atgatattaa aatgattcga
                                                                       180
tcacagtcag gtattcatgg ttttgaaccc tcatcaagtg atgttgctgc catttatgat
                                                                       240
gctgatctat ttctttatca ttcgcacaca ctagaagctt gggcgagacg tttggaacct
                                                                       300
agttttgcatc actctaaagt atctgtaatt gaagcttcaa aaggtatgac tttggataaa
                                                                       360
gttcatggct tagaagatgt agaggcagaa aaaggagtag atgagtcaac cttgtatgac
                                                                       420
cctcacactt ggaatgaccc tgtaaaagta tctgaggaag cacaactcat cgctacacaa
                                                                       480
ttagctaaaa aggatcctaa aaacgctaag gtttatcaaa aaaatgctga tcaatttagt
                                                                       540
gacaaggcaa tggctattgc agagaagtat aagccaaaat ttaaagctgc aaagtctaaa
                                                                       600
tactttgtga cttcacatac agcattctca tacttagcta agcgatacgg attgactcag
                                                                       660
ttaggtattg caggtgtctc aaccgagcaa gaacctagtg ctaaaaaatt agccgaaatt
                                                                       720
caggagtttg tgaaaacata taaggttaag actatttttg ttgaagaagg agtctcacct
                                                                       780
aaattagctc aagcagtagc ttcagctact cgagttaaaa ttgcaagttt aagtccttta
                                                                       840
raagcagttc ccaaaaacaa taaagattac ttagaaaatt tggaaactaa tcttaaggta
                                                                       900
cttgtcaaat cgttaaatca atag
                                                                       924
<210> 40
<211> 307
<212> PRT
<213> Streptococcus agalactiae
<220>
<221> VARIANT
<222> (1)...(307)
<223> Xaa = Any Amino Acid
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Leu Ile Cys Leu Cys Ala Cys Thr Lys Gln Ser Gln Gln Lys Asn Gly
                                25
Leu Ser Val Val Thr Ser Phe Tyr Pro Val Tyr Ser Ile Thr Lys Ala
Val Ser Gly Asp Leu Asn Asp Ile Lys Met Ile Arg Ser Gln Ser Gly
                        55
Ile His Gly Phe Glu Pro Ser Ser Ser Asp Val Ala Ala Ile Tyr Asp
                                        75
Ala Asp Leu Phe Leu Tyr His Ser His Thr Leu Glu Ala Trp Ala Arg
                85
                                    90
Arg Leu Glu Pro Ser Leu His His Ser Lys Val Ser Val Ile Glu Ala
```

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100
                                105
                                                     110
Ser Lys Gly Met Thr Leu Asp Lys Val His Gly Leu Glu Asp Val Glu
                            120
Ala Glu Lys Gly Val Asp Glu Ser Thr Leu Tyr Asp Pro His Thr Trp
                        135
                                             140
Asn Asp Pro Val Lys Val Ser Glu Glu Ala Gln Leu Ile Ala Thr Gln
                    150
                                        155
Leu Ala Lys Lys Asp Pro Lys Asn Ala Lys Val Tyr Gln Lys Asn Ala
                165
                                    170
Asp Gln Phe Ser Asp Lys Ala Met Ala Ile Ala Glu Lys Tyr Lys Pro
            180
                                185
                                                     190
Lys Phe Lys Ala Ala Lys Ser Lys Tyr Phe Val Thr Ser His Thr Ala
                            200
Phe Ser Tyr Leu Ala Lys Arg Tyr Gly Leu Thr Gln Leu Gly Ile Ala
                        215
Gly Val Ser Thr Glu Gln Glu Pro Ser Ala Lys Lys Leu Ala Glu Ile
225
                    230
                                        235
Gln Glu Phe Val Lys Thr Tyr Lys Val Lys Thr Ile Phe Val Glu Glu
                245
                                    250
Gly Val Ser Pro Lys Leu Ala Gln Ala Val Ala Ser Ala Thr Arg Val
            260
                                                     270
                                265
Lys Ile Ala Ser Leu Ser Pro Leu Xaa Ala Val Pro Lys Asn Asn Lys
                            280
Asp Tyr Leu Glu Asn Leu Glu Thr Asn Leu Lys Val Leu Val Lys Ser
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                                             300
Leu Asn Gln
305
<210> 41
<211> 1134
<212> DNA
<213> Streptococcus agalactiae
<400> 41
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                                                                       120
aacgaaaaat tacgcttaga taaaagaagt aaattaaata tttcttctcc tqaaqaacct
                                                                       180
caaaatacta ctaaaattaa gaagcttcat tttccaaaga tttcaagacc taagattgaa
                                                                       240
aagaaacaga aaaaagaaaa aatagtcaac agcttagcca aaactaatcg cattagaact
                                                                       300
gcacctatat ttgtagtagc attcctagtc attttagttt ccqttttcct actaactcct
                                                                       360
tttagtaagc aaaaaacaat aacagttagt ggaaatcagc atacacctga tgatattttg
                                                                       420
atagagaaaa cgaatattca aaaaaacgat tatttctttt ctttaatttt taaacataaa
                                                                       480
gctattgaac aacgtttagc tgcagaagat gtatgggtaa aaacagctca gatgacttat
                                                                       540
caatttccca ataagtttca tattcaagtt caagaaaata agattattgc atatgcacat
                                                                       600
                                                                       660
acaaagcaag gatatcaacc tgtcttggaa actggaaaaa aggctgatcc tgtaaatagt
tcagagctac caaagcactt cttaacaatt aaccttgata aggaagatag tattaagcta
                                                                       720
ttaattaaag atttaaaggc tttagaccct gatttaataa gtgagattca ggtgataagt
                                                                       780
ttagctgatt ctaaaacgac acctgacctc ctgctgttag atatgcacga tggaaatagt
                                                                       840
                                                                       900
attagaatac cattatctaa atttaaagaa agacttcctt tttacaaaca aattaagaag
aaccttaagg aaccttctat tgttgatatg gaagtgggag tttacacaac aacaaatacc
                                                                       960
attgaatcaa cccctgttaa agcagaagat acaaaaaata aatcaactga taaaacacaa
                                                                      1020
acacaaaatg gtcaggttgc ggaaaatagt caaggacaaa caaataactc aaatactaat
                                                                      1080
caacaaggac aacagatagc aacagagcag gcacctaacc ctcaaaatgt taat
                                                                      1134
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<212> PRT
<213> Streptococcus agalactiae
Met Pro Lys Lys Ser Asp Thr Pro Glu Lys Glu Glu Val Val Leu
Thr Glu Trp Gln Lys Arg Asn Leu Glu Phe Leu Lys Lys Arg Lys Glu
                               25
Asp Glu Glu Glu Gln Lys Arg Ile Asn Glu Lys Leu Arg Leu Asp Lys
                            40
Arg Ser Lys Leu Asn Ile Ser Ser Pro Glu Glu Pro Gln Asn Thr Thr
                        55
                                            60
Lys Ile Lys Lys Leu His Phe Pro Lys Ile Ser Arg Pro Lys Ile Glu
Lys Lys Gln Lys Lys Glu Lys Ile Val Asn Ser Leu Ala Lys Thr Asn
                85
                                    90
Arg Ile Arg Thr Ala Pro Ile Phe Val Val Ala Phe Leu Val Ile Leu
                                105
Val Ser Val Phe Leu Leu Thr Pro Phe Ser Lys Gln Lys Thr Ile Thr
                           120
Val Ser Gly Asn Gln His Thr Pro Asp Asp Ile Leu Ile Glu Lys Thr
                       135
                                            140
Asn Ile Gln Lys Asn Asp Tyr Phe Phe Ser Leu Ile Phe Lys His Lys
                   150
                                       155
Ala Ile Glu Gln Arg Leu Ala Ala Glu Asp Val Trp Val Lys Thr Ala
               165
                                   170
Gln Met Thr Tyr Gln Phe Pro Asn Lys Phe His Ile Gln Val Gln Glu
            180
                                185
Asn Lys Ile Ile Ala Tyr Ala His Thr Lys Gln Gly Tyr Gln Pro Val
                            200
Leu Glu Thr Gly Lys Lys Ala Asp Pro Val Asn Ser Ser Glu Leu Pro
                       215
                                           220
Lys His Phe Leu Thr Ile Asn Leu Asp Lys Glu Asp Ser Ile Lys Leu
                   230
                                       235
Leu Ile Lys Asp Leu Lys Ala Leu Asp Pro Asp Leu Ile Ser Glu Ile
               245
                                   250
Gln Val Ile Ser Leu Ala Asp Ser Lys Thr Thr Pro Asp Leu Leu Leu
                               265
Leu Asp Met His Asp Gly Asn Ser Ile Arg Ile Pro Leu Ser Lys Phe
                           280
Lys Glu Arg Leu Pro Phe Tyr Lys Gln Ile Lys Lys Asn Leu Lys Glu
                        295
                                            300
Pro Ser Ile Val Asp Met Glu Val Gly Val Tyr Thr Thr Thr Asn Thr
                    310
                                        315
Ile Glu Ser Thr Pro Val Lys Ala Glu Asp Thr Lys Asn Lys Ser Thr
                325
                                    330
Asp Lys Thr Gln Thr Gln Asn Gly Gln Val Ala Glu Asn Ser Gln Gly
                               345
Gln Thr Asn Asn Ser Asn Thr Asn Gln Gln Gly Gln Gln Ile Ala Thr
                            360
Glu Gln Ala Pro Asn Pro Gln Asn Val Asn
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